Factors influencing *Prymnesium parvum* population dynamics during bloom initiation: Results 1 2 from in-lake mesocosm experiments 3 Daniel L. Roelke*¹, Leslie Schwierzke², Bryan W. Brooks³, James P. Grover⁴, Reagan M. Errera², 4 Theodore W. Valenti Jr.⁵, James L. Pinckney⁶ 5 6 ¹Associate Professor, corresponding author 7 8 Departments of Wildlife and Fisheries Sciences, and Oceanography 9 Texas A&M University 10 **2258 TAMUS** 11 College Station, TX 77843-2258, USA Phone: (409) 845-0169; Fax: (409) 845-4096; E-mail: droelke@tamu.edu 12 13 14 ²Research Associate 15 Department of Wildlife and Fisheries Sciences 16 Texas A&M University 17 ³Associate Professor 18 19 Department of Environmental Science, and Center for Reservoir and Aquatic Systems Research 20 **Baylor University** 21 22 ⁴Professor 23 Department of Biology, and Program in Environmental and Earth Sciences 24 University of Texas at Arlington 25 26 ⁵Research Associate 27 Department of Environmental Science, and Center for Reservoir and Aquatic Systems Research 28 **Baylor University** 29 30 ⁶Associate Professor Marine Science Program and Department of Biological Sciences 31 32 University of South Carolina 33 34 35 Running head: 36 P. parvum bloom formation 37 38

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Abstract

The alga <i>Prymnesium parvum</i> forms large fish-killing blooms in many Texas (USA) lakes.
In some of these lakes, however, P. parvum occurs but does not develop blooms. In this study,
we investigated factors that may influence bloom initiation by conducting a series of in-lake
experiments involving mixing of waters from L. Whitney, which has a history of <i>P. parvum</i>
blooms, with waters from L. Waco where no blooms have occurred. In all experiments, the
addition of L. Waco waters resulted in a poorer performance of <i>P. parvum</i> . Various experimental
treatments and field data show that differences in grazing, pathogens, nutrients, and salts between
the two lakes were not likely factors that contributed to this observation. Industrial and
agricultural contaminants, allelochemicals and algicidal chemicals were not measured as a part of
this research. However, anthropogenic contaminants other than nutrients were not observed at
levels exceeding water quality standards in L. Waco in recent years. On the other hand, nuisance
cyanobacteria are common in L. Waco, where Microcystis sp. and Anabaena sp. were abundant
during the initiation of our experiments, both taxa are known to produce chemicals with
allelopathic properties. In addition, the emergent field of algal-heterotrophic bacteria interactions
suggests that chemicals produced by heterotrophic bacteria should not be overlooked. Further
research focusing on the chemical interactions between cyanobacteria and <i>P. parvum</i> , as well as
the potential role of algicidal bacteria, in the initiation of <i>P. parvum</i> blooms is necessary, as it
may be important to the management of these blooms.

Keywords:

plankton, inorganic nutrients, salinity, grazing, harmful algal blooms, microcystins, prymnesins,

allelopathy, cyanobacteria

Introduction

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Prymnesium parvum is a haptophyte alga that occurs worldwide, tolerates large variations in temperature and salinity, and sometimes forms blooms that result in large fish kills (Edvardsen and Paasche 1998, Lundholm and Moestrup 2006, Baker et al. 2007). In Texas (USA), the incidence of *P. parvum* blooms has increased dramatically since 2001, where blooms are now observed in over 19 lakes found along five river basins. During blooms, surface waters take on a golden color with P. parvum cell densities typically exceeding 10×10^6 cells L⁻¹. In addition, fish kills occurring with these blooms involve many species, where total mortalities number in the tens of millions (TPWD 2003, Roelke et al. 2007). Changes in the physicochemical environment that led to the increase in Texas P. parvum blooms are unknown, but might include eutrophication and salinization. For example, blooms in Europe, the Middle East, and Asia have all occurred in aquatic systems that were eutrophic and brackish (Krasnotshchek and Abramowitsch 1971, Holdway et al. 1978, Rijn and Shilo 1989, Kaartvedt et al. 1991, Guo et al. 1996, Amsinck et al. 2005). P. parvum blooms in Texas are partly consistent with these observations because they appear mostly in lakes that attain salinities 2 to 4 psu (practical salinity units) during low precipitation years (TPWD 2003), and these lakes may have experienced increased non-point source nutrient loading because of aging septic systems, point source discharges and expanded shoreline development. However, the role of nutrients is complex. In laboratory and field experiments the toxicity to fish from chemicals produced by P. parvum, which also act as allelochemicals and are an important factor leading to bloom initiation, was greater when cells were nutrient-limited (Uronen et al. 2005, Roelke et al. 2007, Errera et al. 2008). In addition, P. parvum is sensitive to pulses of nutrients where high

doses inhibited bloom formation (Barkoh et al. 2003, Grover et al. 2007, Kurten et al. 2007).

Therefore, the temporal variation in nutrient availability is likely more important than a system's trophic state.

In general, factors leading to the formation of harmful algal blooms (HABs) are many and diverse (see Paerl 1988, Roelke and Buyukates 2001). In regards to *P. parvum*, bloom-initiating processes may include production of chemicals toxic to grazers (Granéli and Johansson 2003, Tillmann 2003, Barreiro et al. 2005, Calliari and Tiselius 2005, Roelke et al. 2007, Brooks et al. *Accepted, this issue*), use of alternative energy and nutrient sources through mixotrophy (Nygaard and Tobiesen 1993, Skovgaard and Hansen 2003), suppression of competitors through allelopathy (Fistarol et al. 2003, 2005, Granéli and Johansson 2003, Roelke et al. 2007, Errera et al. 2008, Grover et al. *Accepted, this issue*), and resistance to the allelopathic effects of other algae (Suikkanen et al. 2004, Tillmann et al. 2007). These factors are not likely mutually exclusive. Additional factors that seem to influence the growth of other HABs, which might also influence *P. parvum* population dynamics, include the production of beneficial or deleterious chemicals by various bacteria taxa (Kodama et al. 2006, Salomon and Imai 2006) and the pathogenic effects of some virus (Salomon and Imai 2006, Schwierzke et al. *In Review, this issue*).

Interestingly, *P. parvum* occurs in some Texas lakes where blooms do not form. At present, it is unclear why blooms do not develop in these lakes while occurring in neighboring reservoirs. As previously mentioned, factors influencing bloom formation might involve various aspects within the water environment ranging from zooplankton grazers to dissolved chemicals. The overall objective of this research was to determine the relative importance of these factors, or narrow down the number of potential influences on *P. parvum* bloom formation. To achieve this

goal, we conducted in-lake mesocosm experiments involving the mixing of waters from a lake frequented by *P. parvum* blooms with waters from a lake unaffected by blooms. The experimental treatments included manipulations of initial grazer and competitor population densities and community composition, along with manipulations of various ions and dissolved chemicals. Our experiments were conducted during the time of year when blooms start to develop in many Texas lakes.

Materials and Methods

L. Whitney is a reservoir on the Brazos River, Texas, USA, constructed in 1951. The lake receives drainage from an area of 42,107 km², has a capacity of 4.68 x 108 m³, a surface area of 95 km², and a shoreline of 362 km (Bailes and Hudson 1982). River discharge in this developed watershed is flashy, with peak flows typically occurring in the late spring and early summer, lasting several days, which can result in hydraulic flushing peaks of ~0.25 d¹. Discharge during the late summer, fall and winter months is low and sometimes undetectable. We chose an area further south in the lake for our study site (32°52.50, 97°23.10) where historically high *P. parvum* population densities have been observed during the late-fall through early-spring months, typically November through March (Glass, personal communication). L. Waco, a nearby reservoir from which waters were transported for these experiments, is located within the Bosque watershed. Relative to L. Whitney, L. Waco is smaller in area and capacity (78.7 km², 1.88 x 108 m³) and drains less land area (4,325 km²), but it is similar in geomorphology and hydrology, where river discharge in the late spring and early summer can result in hydraulic flushing peaks of ~0.33 d¹ (Dowell 1972, Lind and Barcena 2003).

We performed three in-lake experiments during a 5-week period in fall 2006, each lasting 7 days. Previous experiments conducted in a neighboring system observed significant phytoplankton and zooplankton responses to treatments within a 7-day period (Roelke et al. 2007, Errera et al. 2008). In experiments employing smaller volume containers, however, experimental artifacts were observed beyond the 7-day period (Errera et al. 2008). Consequently, we selected a 7-day duration for our experiments. Over short periods, the emerging plankton composition resulting from competition and other foodweb interactions can be sensitive to the initial community composition (Roelke et al. 2003, Roelke and Eldridge *In Press*). Since the community composition shifts during the time of year when blooms are forming in Texas lakes, possibly influencing the outcome of our treatments, we repeated three consecutive experiments, initiated October 3rd, 17th, and 31st.

In this study, we utilized 36 transparent 25-L polycarbonate carboys during each experiment, filled to a final volume of 24.5 L. Air in the headspace of each carboy allowed neutral buoyancy. The carboys were suspended in the near-surface water from anchored floatation platforms that allowed free movement with wave turbulence, keeping the carboys well mixed. To simulate the natural light environment, the carboys were covered with a neutral density screen, reducing surface light by ~55%. Lake secchi depths are typically ~1 m at this time of year, which translates to a light extinction coefficient of ~1.7 m⁻¹ (see Wetzel 2001). A 55% reduction in surface light would then occur at ~0.5 m, the depth from which water was collected for experiment initiation. Measurements of pH taken at initiation and termination of the 7-day experiments suggested that CO₂ did not limit primary productivity in the sealed carboys.

Lake water used to initiate these experiments was collected from locations within L. Whitney and L. Waco away from the shoreline, where *P. parvum* occurred at the time. Profiles of temperature and salinity suggested that surface waters from both reservoirs were well mixed at the time of collection. Previous studies have identified that pH levels can influence the potency of prymnesins, toxins produced by *P. parvum* (Ulitzur and Shilo 1966, Shilo 1981). However, pH values between the lakes differed by only 0.25, with a mean value of 8.21. This slight variation did not cause differential potency effects between the lake waters.

Each experiment comprised 12 treatments that were conducted in triplicate. Five of the treatments represented a gradient of combined, unfiltered waters from L. Whitney and L. Waco at proportions of 100:0, 75:25, 50:50, 25:75 and 0:100 (Whitney:Waco). In these treatments, our goal was to assess the combined effect of grazers, competitors, pathogens and dissolved chemicals from L. Waco on population dynamics of *P. parvum*.

Another of our goals was to differentiate between the combined effects of grazers and competitors from L. Waco, from the combined effects of virus pathogens and dissolved chemicals from L. Waco on the population dynamics of *P. parvum*. To accomplish this, two additional treatments were performed. The first included a 50:50 mixed proportion of unfiltered L. Whitney water and 0.45 µm filtered (membrane filter) L. Waco water; the second, a 50:50 mixed proportion of unfiltered and 0.45 µm filtered L. Whitney waters. We assumed the filtered waters primarily contained dissolved constituents, viral particles, and bacteria <0.45 µm in cell size. Both treatments had the same initial population densities of phytoplankton and zooplankton, at 50% of natural abundances in L. Whitney, allowing us to account for potential effects of varied initial conditions on emergent community composition (Roelke et al. 2003, Roelke and Eldridge *In Press*).

Additional goals of our research were to investigate the roles of nutrients and salinity on P. parvum population dynamics when L. Waco and L. Whitney waters were mixed. To achieve this, a final five treatments were added. The first three involved mixed proportions of 100:0. 50:50 and 0:100 of unfiltered waters from L. Whitney and L. Waco with additions of inorganic nutrients, trace metals and vitamins to f/2 concentrations (Guillard and Ryther 1962). In regards to N and P, enrichment of N was to 800 µM-N and enrichment of P was to 40 µM-P (N:P ratio of 20). By enriching these treatments, the chance of growth limitation arising from the scarcity of inorganic nutrients, trace metals and vitamins was negated. The remaining two treatments included a 50:50 mixed proportion of waters with salts added, and a 50:50 mixed proportion of water from L. Whitney and 0.45 µm filtered water from L. Waco with salts added. Salinity levels were adjusted to match those found in L. Whitney. During our in-lake experiments, L. Waco salinity was ~0.14 psu, and L. Whitney salinity averaged 1.73 psu. A defined medium of artificial seawater (Kester 1967) was added to each carboy in order to reach the higher salinity characteristic of L. Whitney. Based on our observations while culturing the Texas strain of P. parvum, where inoculations are periodically performed into media with salinities varying by as much as 3 psu at low salinity ranges, it is not likely that an osmotic-shock due to a sudden shift of ~ 0.7 psu (from 1.73 to ~ 1 psu) influenced the outcome of our 7-day experiments. Response variables measured in our experiments included characterizations of plankton, inorganic nutrients, and ambient toxicity (where toxicity refers to the net affect of all chemicals). Characterizations of plankton included estimates of total phytoplankton biomass and biomasses of higher taxonomic groups, enumerations of *P. parvum* population densities, and total

zooplankton biomass and biomasses of higher taxonomic groups. Characterizations of the

nutrients included measurements of nitrate (NO₃), nitrite (NO₂), ammonium (NH₄), and

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orthophosphate (PO₄). Ambient toxicity was estimated using standardized fish and cladoceran bioassays previously described (Roelke et al. 2007). Initial conditions were characterized by measurements taken just prior to the initiation of each experiment from both source waters.

Response variables were then sampled from each carboy at the end of each 7-day experiment.

Estimates of total phytoplankton biomass and biomasses of higher taxa were determined from phytopigment concentration measurements (Pinckney et al. 1998) and the use of CHEMTAX, a matrix factorization program that enables estimates of taxonomically aggregated assemblage composition (Mackey et al. 1997, Wright et al. 1996). For the CHEMTAX model initiation, cyanobacteria, euglenophytes, chlorophytes, prymnesiophytes, cryptophytes, diatoms and chrysophytes were selected because of their historical prevalence in L. Whitney and L. Waco. For greater detail of the HPLC and CHEMTAX methods followed, see Roelke et al. (2007).

Water column chlorophyll *a* concentrations were also determined using standard fluorometric procedures. Triplicate 50 mL samples were filtered through 47mm GF/F filters per carboy and frozen until analysis. Pigments were extracted with 90% acetone, centrifuged, and analyzed using a fluorometer (APHA 1998).

From each carboy a 100 mL phytoplankton sample was collected and preserved using glutaraldehyde, 5% v/v. Enumeration of *P. parvum* population density was performed using a settling technique (Utermöhl 1958). A subsample of 0.5 to 1.5 mL was settled for 24 h, then 30-40 randomly selected fields of view were counted using an inverted, phase contrast light microscope (400x, Leica Microsystems). Total cells counted per sample averaged between 50-150.

Detailed microscopy was performed on L. Whitney and L. Waco waters used to initiate the experiments, following the same methods described above. The focus of this effort was to determine the proportion of the total phytoplankton biovolume that was represented by specific cyanobacteria taxa, i.e., genera-level characterizations that are not possible using the CHEMTAX model.

Zooplankton samples were collected following two methods. For initial in-lake conditions a Schindler trap (61 μm mesh size) was used to concentrate a 12 L sample to 50 ml. To sample carboys at the termination of each experiment, 10 L were removed and filtered through the cod end portion of a Schindler trap and concentrated to 50 ml. All zooplankton samples were preserved in buffered formalin, 5% v/v. A subsample of 5 to 15 mL was settled for 24 hours, then counted using an inverted, phase contrast light microscope (40x and 200x, Leica Microsystems). For each individual counted, dimensions were measured corresponding to best-fit geometric shapes to estimate biovolume (Wetzel and Likens 1991). For this study zooplankton species were grouped into total copepod adults, copepod nauplii, total rotifers, total cladocerans and total protozoans (ciliate and amoeboid). Our enumeration technique resulted in ~100-150 total individuals counted per sample.

Samples for inorganic nutrients were filtered through GF/F filters, and the filtrates were frozen until analysis. Inorganic nutrient concentrations were determined using autoanalyzer methodology (Armstrong and Sterns 1967, Harwood and Kuhn 1970). For this study, NO₃, NO₂ and NH₄ were summed (DIN).

Ambient acute toxicity to fish was evaluated for initial conditions and from each experimental carboy using standardized 24-hour static toxicity assays with the juvenile fathead minnow (*Pimephales promelas*) model. Sublethal toxicity to a model cladoceran was evaluated

with standardized 10-day static renewal chronic toxicity tests with the *Daphnia magna* model. Toxicity assays followed standardized methods for aquatic toxicology (US EPA 1994, 2002). Samples were collected and transported to the laboratory where toxicity tests were initiated within 24 hours. Ambient samples were diluted using a 0.5 dilution series with reconstituted hard water (RHW), which was performed according to US EPA recommendations (US EPA 2002). For greater detail of the methods followed for these toxicity assays, refer to Brooks et al. (2004) and Dzialowski et al. (2006).

Using SPSS 14, comparisons of the response variables ([day 7 – day 0]/day 0) between experimental treatments involving the five mixed proportions of unfiltered waters from L. Whitney and L. Waco were tested using one-way ANOVA, followed by Bonferroni and Tukey *post hoc* tests. Independent samples t-tests were performed for all other treatment comparisons. These comparisons focused on the differences of primary interest in this study: effects of the source and proportion of waters mixed in the experimental carboys.

Results

In-situ conditions for L. Whitney and L. Waco

From the three sampling dates in L. Whitney (October 3^{rd} , 17^{th} and 31^{st}), *P. parvum* cell densities were 0.72×10^6 , 0.50×10^6 , and 0.83×10^6 cells L⁻¹, well below bloom proportions in Texas waters ($\sim 10 \times 10^6$ cells L⁻¹), and toxicity bioassays confirmed non-toxic in-lake conditions. The state fisheries managers indicated no observations of golden-colored waters and no reports of fish kills at this time (Glass, personal communication). Foam lines were present, however, which typically correspond with the incidence of *P. parvum*. Approximately one month after our inlake experiments golden-colored waters and fish kills were observed. Therefore, we conclude that

these experiments took place during the period of bloom initiation.

have still been no observed P. parvum blooms in L. Waco.

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Cyanobacteria dominated the phytoplankton assemblage (50% of the biovolume, almost all *Phormidium* sp.; Table 1) during the October 3rd sampling of L. Whitney. During the later two October samplings, phytoplankton composition was more diverse, with co-dominance of chlorophytes, chrysophytes, cyanobacteria and prymnesiophytes, and sub-dominance of euglenophytes and cryptophytes. Diatoms were below detection based on CHEMTAX analysis of pigments, and cyanobacteria known to be harmful were not observed in microscopy (Table 1). Total phytoplankton biomass fluctuated during the three lake samplings, with concentrations of 17.9, 25.9 and 19.8 μ g-chl $a L^{-1}$. Total zooplankton biovolume was similar for the first two samplings in L. Whitney, but then increased ~10-fold over the study period. On October 3rd and 17th biovolumes were 3.0 x 10⁷ and 1.3 x 10⁷ μm³ L⁻¹, but on October 31st total biovolume was 13.8 x 10⁷ μm³ L⁻¹. Rotifers accounted for 86% of the total biovolume on October 3rd (101 individuals L⁻¹), but by October 17th rotifers declined to 37% (16 individuals L⁻¹) and copepod nauplii comprised 63% (10 individuals L⁻¹) of the total biovolume. By October 31st rotifers became subdominant (15%, 70 individuals L⁻¹), and copepod nauplii (44%, 103 individuals L⁻¹) and copepod adults (34%, 7 individuals L⁻¹) were co-dominant. Cladocerans, and ciliated and amoeboid protozoa were not observed. L. Waco was sampled on the same dates as L. Whitney, and *P. parvum* cell densities were very similar at 0.31 x 10⁶, 0.37 x 10⁶, and 0.62 x 10⁶ cells L⁻¹. Consistent with this lake's history, there were no signs of *P. parvum* blooms during the period of our experiments. To date, there

During October, total phytoplankton biomass in L. Waco was similar to levels found in L. Whitney: 17.4, 19.6 and 21.5 μg-chl *a* L⁻¹ for the three sampling dates in L. Waco. However, cyanobacteria were much more prevalent compared to L. Whitney, accounting for 77%, 69% and 51% of the total phytoplankton biovolume for the three sampling dates. In addition, potentially toxigenic species were observed in our microscopy, where *Microcystis* sp. and *Anabaena* sp. were dominant (Table 1).

Zooplankton varied in L. Waco during October, with total biovolumes of 10.7×10^7 , 1.3×10^7 and 10.0×10^7 µm³ L⁻¹ for the three sampling dates. Similar to L. Whitney, initial rotifer dominance gave way to co-dominance with copepod nauplii. On October 3rd rotifers accounted for 98% (114 individuals L⁻¹) of the total zooplankton biovolume, but on October 17th and 31st, rotifers accounted for 41% and 53% of the total biovolume (25 and 93 individuals L⁻¹). Copepod nauplii shared dominance at 57% and 37% of the total biovolume (11 and 46 individuals L⁻¹) during the latter two sampling dates. Similar to L. Whitney, cladoceran and ciliated and amoeboid protozoa population densities were very low in L. Waco during this time.

Inorganic nutrient concentrations were similar between the two lakes. In L. Whitney, average DIN and PO_4 concentrations (from the three lake samples collected at the start of each experiment) were 3.8 μ M-N and 0.10 μ M-P, and in L. Waco they were 2.78 μ M-N and 0.22 μ M-P. The ratio of N:P varied, however; L. Whitney N:P was ~38, while the L. Waco N:P was ~13.

Experimental results

The accumulation of *P. parvum* populations in the experimental carboys varied according to the proportions of waters mixed from L. Whitney and L. Waco. In all three experiments, *P.*

parvum increased 3- to 7-fold in carboys with only unfiltered L. Whitney water, while appreciable increases were not observed in the carboys with only unfiltered L. Waco water (Fig. 1). Furthermore, a dose-response was observed in all three experiments where the accumulation of P. parvum populations diminished with an increasing proportion of unfiltered L. Waco water. Distinct groupings along this mixing gradient were statistically significant (p < 0.05). Low accumulation of P. parvum was also observed in the carboys with filtered waters from L. Waco in all three experiments (Fig. 1). Statistically significant reductions in fish survival and cladoceran reproduction responses were never observed, including in the carboys with greater proportions of L. Whitney water where P. parvum population densities reached near-bloom proportions.

For phytoplankton biomass, mixing of unfiltered waters from L. Whitney and L. Waco did not produce a consistent trend as seen with *P. parvum* (Fig. 2). Reductions in chlorophyll *a* from initial conditions occurred, and the decrease was greater in carboys containing larger proportions of L. Whitney water in the first two experiments, but the opposite was observed for the third experiment.

The phytoplankton response in the carboys containing filtered L. Waco waters was consistent with the trends observed for P. parvum, where in all three experiments performance of the total phytoplankton assemblage was poorer in carboys with L. Waco waters, with significant differences (p < 0.05) for the first and third experiments (Fig. 2). Not all taxa were affected similarly. For example, cyanobacteria, euglenophytes and cryptophytes showed significantly (p < 0.05) poorer performance with the addition of filtered water from L. Waco. A poorer performance of chlorophytes was also observed, but only significant (p < 0.05) in the third experiment. Chrysophytes showed no significant difference between these treatments (Table 2).

Diatoms remained below the detection threshold for pigment analysis with the CHEMTAX model.

Zooplankton biovolume increased from initial conditions in all carboys. In the first two experiments the growth of zooplankton biovolume was significantly greater in carboys with a larger proportion of L. Waco waters (p < 0.05, Fig. 3). Growth of copepod nauplii into adult forms was pronounced in the second and third experiments. The accumulation of zooplankton biovolume in the treatments involving filtered waters from L. Waco was not significantly different from the treatments involving filtered waters from L. Whitney (p > 0.05, Fig. 3).

The addition of nutrients showed varying results. Enrichment did not change the deleterious effect of filtered L. Waco waters on P. parvum populations (p < 0.05, Fig. 4). On the other hand, nutrient additions changed the phytoplankton response to addition of filtered L. Waco waters, but the effect was not consistent between experiments. During the first and third experiments, there was no significant difference in phytoplankton growth between enriched carboys with filtered waters from L. Whitney and L. Waco. During the second experiment, however, phytoplankton growth was significantly greater in the enriched carboys with filtered L. Whitney water. Shifts in assemblage composition between the higher taxonomic groups were minor, except in carboys with 50% filtered L. Whitney waters where euglenophytes dropped below the detection level of pigment analysis based on the CHEMTAX model. Accumulation of zooplankton biovolume was much greater with nutrient additions, primarily due to the population growth of rotifers. Differences between filtered water from L. Whitney and from L. Waco, however, were not observed in any of the three experiments.

Salt additions to mixtures of L. Whitney and L. Waco waters, bringing salinity levels up

to those observed in L. Whitney (from \sim 1 to 1.7 psu), did not alter results in any experiment. That is, no significant differences (p < 0.05) were found between L. Whitney and L. Waco waters (50:50) without salt added compared to salt additions for *P. parvum*, chlorophyll a, or zooplankton biovolume (Fig. 5). Similarly, no significant differences (p < 0.05) were found between L. Whitney and filtered L. Waco waters (50:50) without salt added compared to salt additions for these same parameters (Fig. 6). During the first experiment, notable increases in rotifers were observed with the salt additions, but they were highly variable between replicates.

L. Whitney and L. Waco are similar in their trophic state, and this resulted in comparable nutrient concentrations among various treatments. Excluding the carboys with enrichment, nutrients ranged from 0.9 to 1.1 μ M-DIN and 0.13 to 0.15 μ M-PO₄ at the termination of all three experiments.

Discussion

Waters from L. Waco, when mixed with waters from L. Whitney, had a deleterious impact on *P. parvum* in these experiments. Zooplankton prospered in all treatments, with rapid growth of parthenogenetic rotifers and the development of copepod nauplii into adults. Undoubtedly, grazing pressure increased during the 7 days of each experiment. It was not the zooplankton from L. Waco, however, that led to the poorer performance of *P. parvum* because both unfiltered and filtered waters from L. Waco produced the same deleterious response. In addition, accumulation of zooplankton biomass differed significantly between mixing treatments only in the first experiment, where copepod adults increased in biomass. Remaining potential causes for poor performance of *P. parvum* in L. Waco waters then, were some small pathogen passing

through a 0.45 µm filter, possibly a virus, or dissolved chemicals, such as a nutrients, salts, contaminants, allelochemicals or algicides.

The role of pathogens in phytoplankton population dynamics is becoming clearer with advances in technologies enhancing study in this area. Viruses directly influence other harmful algal species (Brussaard 2004, Salomon and Imai 2006), and there is evidence that viruses influence *P. parvum* population dynamics during later stages of blooms (Schwierzke et al. *In Review, this issue*). However, pathogens tend to be species-specific and do not affect the entire phytoplankton assemblage (Tomaru et al. 2004, Salomon and Imai 2006). In all three of our experiments, additions of filtered L. Waco waters resulted in poorer performance of cyanobacteria, euglenophytes and cryptophytes, which led to an overall poorer performance of the entire L. Whitney phytoplankton assemblage (based on chlorophyll *a*). In addition, signs of pathogens, such as lysed cells, were not observed during microscopy. These lines of evidence indicate that the causative factor in L. Waco waters reducing performance of *P. parvum* was not likely a pathogen.

A more generalized factor that could influence *P. parvum* and the phytoplankton assemblage would be nutrients. However, based on half-saturation coefficients for reproductive growth of many phytoplankton species common to lakes (Grover 1989, Grover et al. 1999), nutrient concentrations would not have been a growth-limiting factor for many species, especially at the start of these experiments. For a Texas strain of *P. parvum*, half-saturation constants for N- and P-limited reproductive growth were estimated to be 0.02 µM or lower (Baker 2007, Baker et al. *Accepted*), indicating that reproductive growth rates would be near maximum during these experiments. More conclusively, the same deleterious effect of filtered L. Waco water on *P*.

parvum was observed in carboys with nutrient enrichment. Further, ammonium levels did not approach those toxic to *P. parvum* or other algae in this study and did not vary appreciably among treatments. For these reasons, it is not likely that differences in nutrient concentrations between L. Whitney and L. Waco led to the poorer performance of *P. parvum*.

For bulk phytoplankton, nutrient limitation would be expected to begin when concentrations drop below 1.0 µM-N and 0.1 µM-P (Reynolds 2006). Interestingly, in the first and third experiments nutrient additions masked the deleterious effect of L. Waco waters when considering the whole phytoplankton community (based on chlorophyll *a*). While major shifts in the aggregated taxonomic groups were not observed, it may be that species shifts within each of the aggregated groups occurred. For example, species resistant to the deleterious effect of L. Waco waters, which could have higher nutrient requirements, might have prospered with the addition of nutrients while sensitive species declined.

Another generalized factor that could influence P. parvum and the phytoplankton assemblage is salinity. For P. parvum, relationships between growth rate and salinity are unimodal or increasing, depending on the strain (Larsen and Bryant 1998). A strain of P. parvum from Texas showed a unimodal relationship with an optimum at 22 psu (Baker et al. 2007). The salinity at the time of our experiments was much lower than that, differing by \sim 1.3 psu between the two reservoirs, and a 50:50 mix of these waters yielded a salinity of \sim 1 psu. While this salinity is low, it is sufficient to support growth of P. parvum when other conditions are favorable (Baker 2007, Baker et al., Accepted).

Applying the relationship between growth and salinity described by Baker (2007), the specific growth rates of *P. parvum* would have been \sim 0.3 and \sim 0.15 d⁻¹ under conditions of \sim 1.7

and ~1 psu. In the treatments where filtered L. Waco waters were mixed with L. Whitney waters and salts were not added (i.e., salinity was not maintained at 1.73), the difference in salinity could have contributed to the differences observed in *P. parvum* population densities. However, when salinity was maintained with the addition of filtered L. Waco waters, the same deleterious effect of filtered L. Waco waters on *P. parvum* was observed, as it was for total phytoplankton.

Therefore, the difference in salt content between the two lakes did not likely cause the negative impact of L. Waco waters on *P. parvum* or the phytoplankton assemblage. Our conclusion is not that salinity has no influence on the incidence of *P. parvum* blooms, as observations suggest otherwise (Krasnotshchek and Abramowitsch 1971, Holdway et al. 1978, Rijn and Shilo 1989, Kaartvedt et al. 1991, Guo et al. 1996, Amsinck et al. 2005). Instead, this indicates that another factor, aside from salinity, played a more significant role in these experiments.

Industrial and agricultural contaminants would also act as generalized factors potentially influencing the entire phytoplankton assemblage. The recent history of L. Waco shows a wide range of contaminants being present, but of the many parameters routinely measured by the Texas Commission on Environmental Quality, only NO₃ was of concern. Parameters not of concern included a suite of heavy metals, polychlorinated biphenyls, organic contaminants (including herbicides), fecal coliform bacteria, suspended sediments, NH₄ and PO₄. Overall, the lake was found to be fully supporting of aquatic life, including phytoplankton (TCEQ 2004). While total phytoplankton is not threatened by present contaminants, individual species might be. Species-specific affects of contaminants have not been studied for L. Waco or L. Whitney, and are well beyond the scope of this research. However, a negative effect on *P. parvum* and major components of the phytoplankton assemblage (cyanobacteria, euglenophytes and

cryptophytes) from contaminants in L. Waco during these experiments is unlikely. On-going monitoring in L. Waco shows accumulation of phytoplankton biomass with diverse taxa and a sustained, but generally small, *P. parvum* population (Roelke, unpublished). If contaminants had an affect similar to the responses measured in our experiments, cyanobacteria, euglenophytes, cryptophytes and *P. parvum* populations would occur at much lower densities then currently observed in L. Waco.

Allelopathy is another generalized factor potentially affecting *P. parvum* and other phytoplankton. *P. parvum* is capable of producing chemicals that suppress the growth of competing phytoplankton (Fistarol et al. 2003, 2005, Granéli and Johansson 2003, Roelke et al. 2007, Errera et al. 2008), as can other harmful species, including *Microcystis* sp. and *Anabaena* sp. (Pflugmacher 2002, Legrand et al. 2003). During our experiments, both of these latter taxa were abundant in waters collected from L. Waco (possibly due to the relatively low N:P) and were absent in waters collected from L. Whitney. Furthermore, in 2006, microcystin-LR was detected by ELISA at levels between 590 and 1090 ng L⁻¹ in L. Waco (Brooks, unpublished), although production of this toxin is not necessary to cause an allelopathic affect (see Sukenik et al. 2002, Beresovsky et al. 2006). Dissolved allelochemicals would have passed through filtration and been present in treatments receiving filtered L. Waco water. Thus, allelopathy could explain why performances of major taxonomic groups in L. Whitney declined with the addition of L. Waco water.

How might *P. parvum* respond to allelochemicals from cyanobacteria? Previous research using a European strain of *P. parvum* indicated a resistance to allelochemicals in studies performed at 6.8 psu (Suikkanen et al. 2004). However, the Texas and European strains have

different photopigments (Errera 2006) and may also vary in vulnerabilities to allelochemicals. The possibility of compounding chemical interactions exists as well. The higher ambient pH of Texas lakes (Roelke, unpublished), relative to the Baltic Sea, may influence the lipophilicity, bioavailability and toxicity of allelochemicals. So, it may be that *P. parvum* populations in L. Whitney are sensitive to cyanobacterial allelochemicals. In addition, a modeling study of L. Granbury, another Texas lake impacted by *P. parvum* blooms, has demonstrated that model behavior better matched in-lake observations of *P. parvum* blooms when allelochemicals from cyanobacteria were considered (Grover et al., *Accepted, this issue*). In addition to cyanobacteria, allelopathy is observed in many other phytoplankton taxa (Granéli and Hansen 2006), therefore allelochemicals from other phytoplankton species in L. Waco cannot be ruled out.

Algicidal chemicals are produced by diverse bacteria and may be another generalized factor potentially affecting *P. parvum* and other phytoplankton (Kodama et al. 2006, Salomon and Imai 2006). The impacts of algicides are also varied. Some broadly affect many phytoplankton taxa, while others are species-specific; in some cases phytoplankton growth is slowed, and in others target populations are obliterated (Doucette et al. 1998, Imai et al. 1998, Yoshinaga et al. 1998, Mayali and Azam 2004). Elucidating the possible role algicidal bacteria played in these experiments was well beyond the scope of this study, but future research should address this possible bloom-suppressing mechanism.

Allelochemicals produced by cyanobacteria, some other phytoplankton taxa, or algicidal chemicals produced by bacteria may be, in part, contributing factors leading to the suppression of *P. parvum* blooms in L. Waco, and perhaps in some other regional lakes as well. For example, in L. Somerville, *P. parvum* is present but does not form blooms. Similar to L. Waco, this lake is

frequently dominated by cyanobacteria (Roelke et al. 2004), and microcystins were detected. In addition, nutrients, dissolved organic carbon, and bacteria concentrations are high. The potential roles of allelochemical-producing phytoplankton and algicidal bacteria as ecosystem-engineering species or in suppressing *P. parvum* blooms in Texas lakes merits further investigation.

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As stated previously, the environmental conditions leading to harmful algal blooms are complex and often species-specific, making it difficult to envision a universal approach to management (Roelke 2000, Roelke and Buyukates 2001, 2002). Our findings, along with those of Grover et al. (Accepted, this issue), suggest that allelopathic phytoplankton or algicidal bacteria might influence P. parvum bloom development. Should follow-on studies confirm this notion, then our findings would have important management implications for *P. parvum* in Texas lakes. For example, a management strategy could focus on the understanding of conditions where P. parvum blooms do not occur, e.g., in the presence of some allelopathic phytoplankton or algicidal bacteria. Manipulating an entire lake to create conditions conducive to a specific microbe may not be feasible, or even wise. However, manipulation of more restricted areas of a lake, timed during the season of bloom initiation, would likely be less detrimental to the overall health of a lake. Targeting specific coves of the dendritic lakes common in this region would be ideal. Ongoing research is investigating the role of coves as bloom initiation "hot spots", possibly due to their longer hydrologic residence time. If true, then a focused, timely effort to promote growth of specific allelopathic phytoplankton or algicidal bacteria, if these are confirmed to suppress P. parvum, in these coves might help to circumvent P. parvum blooms. Even if blooms initiate elsewhere, manipulation in coves may still be advantageous to ward off *P. parvum* blooms, thereby creating a refuge for fish.

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728 **Figure Captions** 729 730 Fig. 1. Population density of *P. parvum* at the start (filled squares) and termination (bars) after 7 days of mixing experiments initiated on October 3rd (A), 17th (B) and 31st (C). Waters 731 represented a gradient of mixing proportions between unfiltered waters from L. Whitney and L. 732 733 Waco; 50% unfiltered L. Whitney waters added to 50% filtered L. Whitney waters; and 50% 734 unfiltered L. Whitney waters added to 50% filtered L. Waco waters. Letter designations 735 represent distinct groupings (p < 0.05). 736 737 Fig. 2. Phytoplankton biomass, as estimated by chlorophyll a concentration, at the start (filled squares) and termination (bars) after 7 days of mixing experiments initiated on October 3rd (A), 738 17th (B) and 31st (C). Waters represented a gradient of mixing proportions between unfiltered 739 740 waters from L. Whitney and L. Waco; 50% unfiltered L. Whitney waters added to 50% filtered 741 L. Whitney waters; and 50% unfiltered L. Whitney waters added to 50% filtered L. Waco waters. Letter designations represent distinct groupings (p < 0.05). 742 743 744 Fig. 3. Average zooplankton biovolume and community composition at the start (filled squares) and termination (stacked bars) after 7 days of mixing experiments initiated on October 3rd (A), 745 17th (B) and 31st (C). Waters represented a gradient of mixing proportions between unfiltered 746 747 waters from L. Whitney and L. Waco; 50% unfiltered L. Whitney waters added to 50% filtered 748 L. Whitney waters; and 50% unfiltered L. Whitney waters added to 50% filtered L. Waco waters. Letter designations represent distinct groupings ($\underline{p} < 0.05$). No letter designations signify no 749

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significant differences were detected..

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752 Fig. 4. Impact of L. Waco waters with the addition of nutrients to population density of P. 753 parvum, phytoplankton biomass, and average zooplankton biovolume and composition at the start (filled squares) and termination (bars) after 7 days for experiments initiated on October 3rd 754 (A), 17^{th} (B) and 31^{st} (C), where the treatments were 50% unfiltered L. Whitney waters added to 755 756 50% filtered L. Whitney waters; and 50% unfiltered L. Whitney waters added to 50% filtered L. 757 Waco waters. In all experiments, L. Waco waters had a deleterious effect on P. parvum 758 population growth (p < 0.5). Phytoplankton response was varied, however, where opposite 759 trends in the first two experiments (A, B) were significant (p < 0.05), and not significant in the 760 third experiment (C). Filtered L. Waco waters had no detectable impact on zooplankton. 761 762 Fig. 5. Impact of L. Waco waters with the addition of salts to match L. Whitney salinities to 763 population density of *P. parvum*, phytoplankton biomass, and average zooplankton biovolume 764 and composition at the start (filled squares) and termination (bars) after 7 days for experiments initiated on October 3rd (A), 17th (B) and 31st (C), where the treatments were 50% unfiltered L. 765 766 Waco waters added to 50% L. Whitney waters; and 50% unfiltered L. Waco waters added to 767 50% L. Whitney waters with salt added. In all experiments, no significant difference (p < 0.5) 768 was detected for P. parvum population growth, or accumulation of phytoplankton and 769 zooplankton biomass. 770 771 Fig. 6. Impact of filtered L. Waco waters with the addition of salts to match L. Whitney 772 salinities to population density of *P. parvum*, phytoplankton biomass, and average zooplankton

biovolume and composition at the start (filled squares) and termination (bars) after 7 days for

experiments initiated on October 3rd (A), 17th (B) and 31st (C), where the treatments were 50% filtered L. Waco waters added to 50% L. Whitney waters; and 50% filtered L. Waco waters added to 50% L. Whitney waters with salt added. In all experiments, no significant difference (*p* < 0.5) was detected for *P. parvum* population growth, or accumulation of phytoplankton and zooplankton biomass.

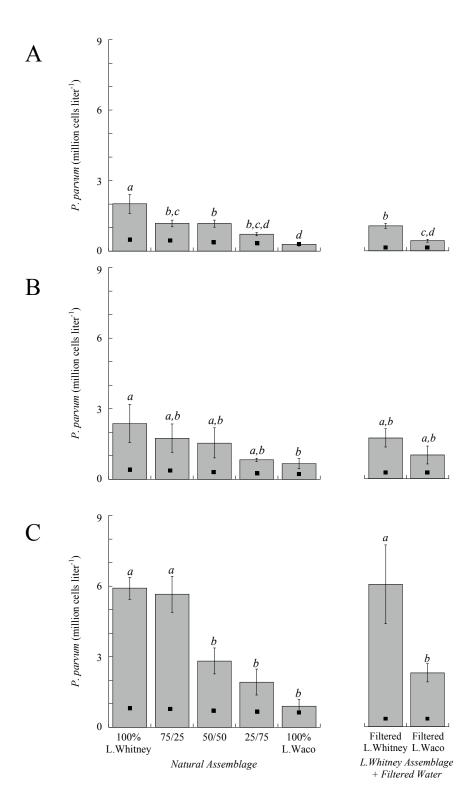


Fig. 1

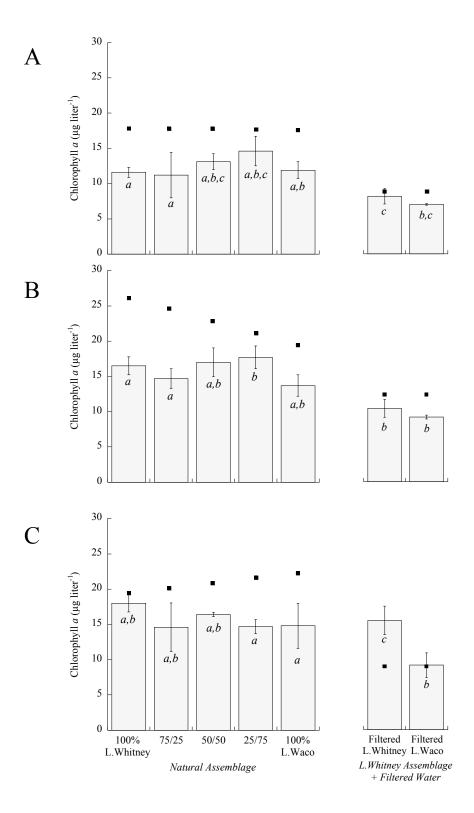


Fig. 2

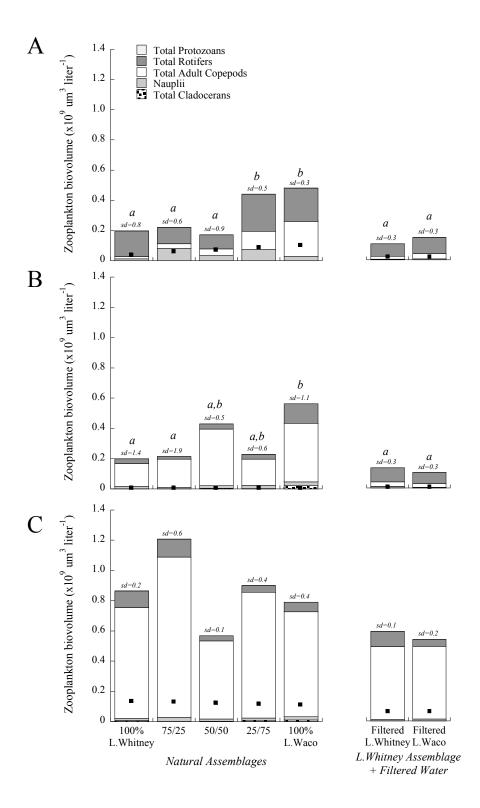


Fig. 3

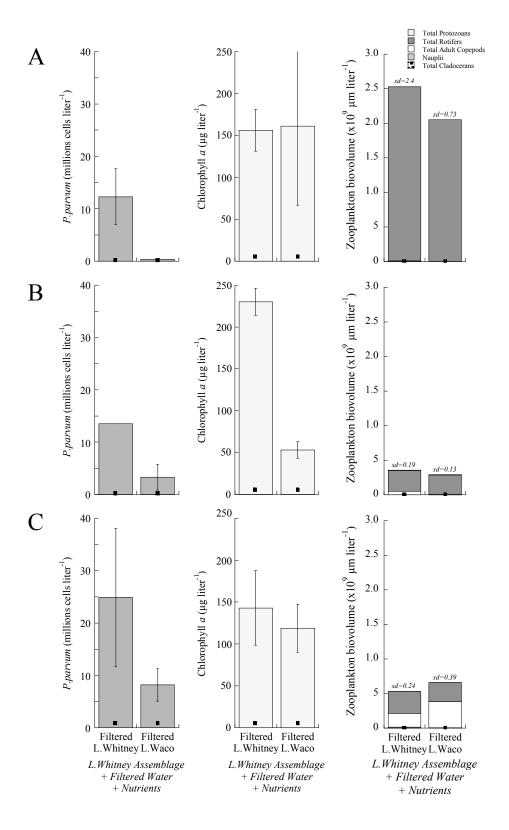


Fig. 4

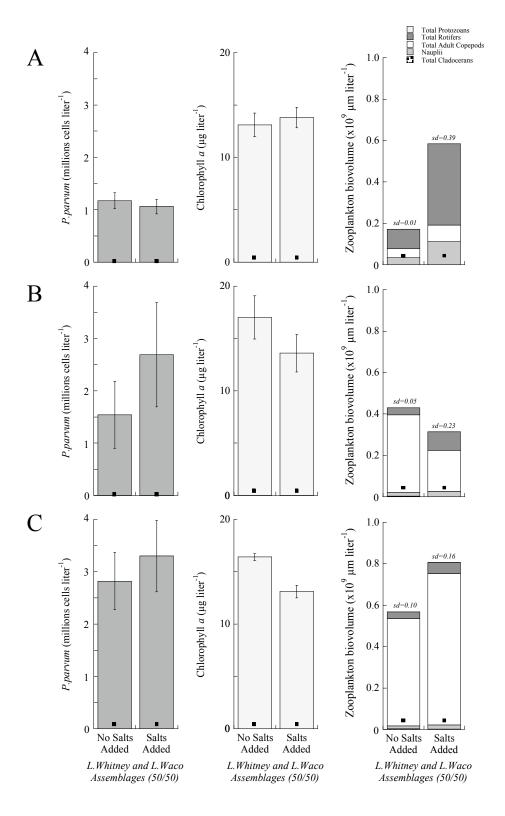


Fig. 5

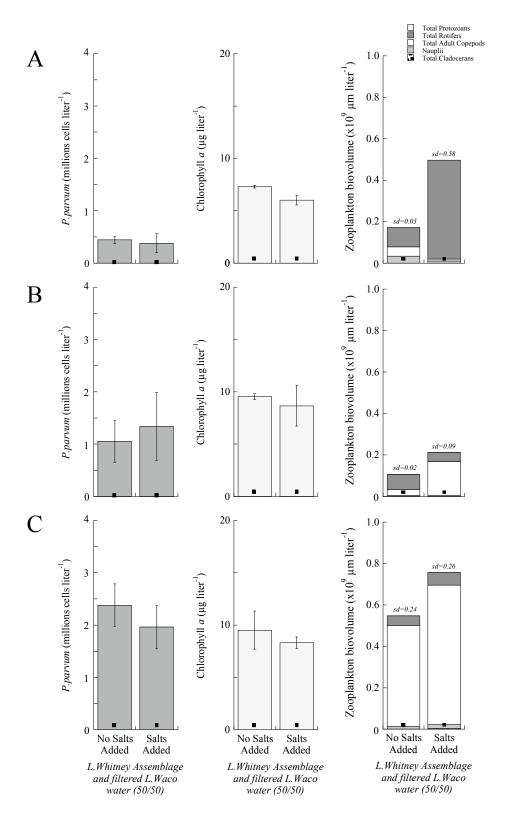


Fig. 6

Table 1. Contributions of cyanobacteria and Prymnesiophytes as percentages of total phytoplankton biovolume at sampling locations in Lakes Whitney and Waco at the start of each experiment.

	Experiment 1 initiated 10-3-06		Experin initiated 1		Experiment 3 initiated 10-31-06	
Taxa	L.Whitney	L.Waco	L.Whitney	L.Waco	L.Whitney	L.Waco
Cyanobacteria	49.5	77.4	18.2	68.7	28.2	51.1
Anabaena sp.	-	18.6	-	_	_	2.4
Calothrix sp.	-	-	_	-	-	5.7
<i>Lyngbya</i> sp.	-	_	-	1.0	-	-
Merismopedia elegans	1.9	2.1	3.8	1.2	1.0	0.8
Merismopedia punctata	0.5	1.5	_	-	-	1.2
Microcystis aeruginosa	-	36.4	_	57.3	8.8	32.9
Phormidium sp.	44.3	7.3	12.9	4.7	15.6	7.3
Raphidiopsis sp.	1.6	2.0	1.0	3.6	1.7	0.9
Synecoccocus sp.	1.2	0.7	0.6	1.0	1.0	-
unknown filament	-	8.7	-	-	-	-
Prymnesiophytes	9.6	1.7	2.8	3.0	6.2	3.5
Other Phytoplankton	40.9	20.9	79.0	28.3	65.6	45.4

Table 2. Average percent change from the initial to the final taxonomic densities (n=3) as measured using CHEMTAX. In the two columns for each experiment results are shown for the treatments where either 50% filtered water from Lake Whitney or 50% filtered water from Lake Waco was added to unfiltered Lake Whitney water.

	Experiment 1		Experiment 2		Experiment 3	
	unfiltered L.Whitney		unfiltered L.Whitney		unfiltered L.Whitney	
	+ filtered	+ filtered	+ filtered	+ filtered	+ filtered	+ filtered
Taxa	L.Whitney	L.Waco	L.Whitney	L.Waco	L.Whitney	L.Waco
Cyanobacteria	29.6	-47*	-14.4	-58.7*	100	-38.2*
Chlorophytes	12.7	15.0	2.7	-9.2	90.6	-4.4*
Euglenophytes	-47.8	-96.8*	11.8	-100*	78.6	-99.5*
Chrysophytes	81.3	65.3	-40.8	-9.7	7.6	26.4
Diatoms	-	-	-	-	-	-
Cryptophytes	-0.2	-53.6*	-19.1	-66.9*	84.4	-30.6*
Prymnesiophytes	57.6	58.1	104	37.7*	221	107*

^{*} significant difference (p < 0.05) between treatments.